REMARKS/ARGUMENTS

Reconsideration of this application and entry of the foregoing amendment are respectfully requested.

The specification has been amended to include the Sequence Listing submitted herewith on separate sheets. Applicants submit that entry of the Sequence Listing does not raise the issue of new matter. The computer readable copy of the Sequence Listing submitted herewith is the same as the attached paper copy of that Listing.

This application is submitted to be in condition for allowance and a Notice to that effect is requested.

Respectfully submitted,

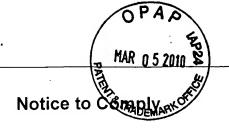
NIXON & VANDERHYE P.C.

By:

Mary J. Wilson

MJW:tat 901 North Glebe Road, 11th Floor Arlington, VA 22203-1808

Telephone: (703) 816-4000 Facsimile: (703) 816-4100



Application No. 10568691	Applicant(s) BAWDEN ET AL.
Examiner Alana M., Harris, Ph.D.	Art Unit 1643

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

1.136(a)).			
The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):			
\boxtimes	1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).		
\boxtimes	2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).		
\boxtimes	3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).		
	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."		
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).		
	6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).		
	7. Other:		
Applicant Must Provide:			
An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".			
\boxtimes	An initial or substitute paper copy of the "Sequence Listing", as well as an amendment specifically directing its entry into the application.		
\boxtimes	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).		
For questions regarding compliance to these requirements, please contact:			
	For Rules Interpretation, call (571) 272-0731 or (571) 272-0951 For CRF Submission Help, call (571) 272-2510 PatentIn Software Program Support Technical Assistance 1-866-217-9197 or 703-305-3028 or 571-272-6845 PatentIn Software is Available At www.USPTO.gov		
PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR REPLY			
	/Alana M. Harris, Ph.D./ Primary Examiner, Art Unit 1643		



<110> Chroma Therapeutics Limited

Bawden, Lindsay J

Bone, Elizabeth A

Drummond, Alan H

· Needham, Lindsey A

- <120> Detection of Histone Modification in Cell-free Nucleosomes
- <130> NRSCP6244818
- <140> PCT/GB2004/003564
- <141> 2004-08-18
- <150> GB 0319376.0
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Val Tyr Leu Ala Ala Val Leu Glu Tyr Leu Thr Ala Glu Ile Leu Glu 50 55 60

Leu Ala Gly Asn Ala Ala Arg Asp Asn Lys Lys Thr Arg Ile Ile Pro 65 70 75 80

Arg His Leu Gln Leu Ala Ile Arg Asn Asp Glu Glu Leu Asn Lys Leu 85 90 95

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Phe Val Asn Asp Ile Phe Glu Arg Ile Ala Gly Glu Ala Ser Arg Leu 65 70 75 80

Ala His Tyr Asn Lys Arg Ser Thr Ile Thr Ser Arg Glu Ile Gln Thr 85 90 95

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